

6 October 2011

Provisional Agenda

- 1 Opening of the meeting
- 2 Nomination of Chairperson, Vice-Chairperson, and Rapporteur
- 3 Adoption of the Provisional Agenda
- 4 General briefing on the working method of the meeting
- 5 Briefing on the background and preparation of updating the WHO/IUCN/WWF guidelines on conservation of medicinal plants (1993)
- 6 Introduction and perspectives of international organizations taking part in the update of the guidelines
 - ♦ WHO
 - ♦ IUCN
 - ♦ WWF
 - ♦ TRAFFIC
 - ♦ FAO
- 7 Review and discussion of the outline of the draft updated guidelines
- 8 Review and discussion of the content of the draft updated guidelines
- 9 General discussion on the future process
- 10 Discussion on recommendations
- 11 Site visit to the relevant research stations in Toyama
- 12 Others, if any
- 13 Closure of the meeting

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10 October 2011

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写真 1 原案提示



写真 2 修正事項討議



写真 3 集合写真

薬用植物を理解

県民公開講座

県民公開講座「意外と知らない身近な薬用植物」は15日、県民会館で開かれ、県民が漢方薬の原料となる薬用植物の現状や作用に理解を深めた。



富山大教員や県職員でつくる「薬用植物の保護に関するWHO会議実行委員会」（委員長・佐竹元吉富山大和漢医薬学総合研究所客員教授）が企画。世界保健機関本部の丸山由紀子科学官、佐竹教授、同研究所の小松かつ子教授、門脇真教授の4人が講演した。

小松教授は、中国産の生薬は採集地の減少や富裕層の増加などで近年、価格が2〜4倍に高騰しているとし、「国内で可能な薬用植物の栽培を推進し、栽培した薬用植物を流通させる仕組みをつくる」ことが必要だと語った。

記事1 北日本新聞 2011年 10月 16日

「薬用植物 保護に力」

WHO担当者ら 知事表敬訪問

きょうから富山 県民会館で専門家会議

15日から富山市の県民会館で薬用植物の保護に関する世界保健機関（WHO）の専門家会議が開かれる。富山県知事（左）とWHOの代表者らと表敬訪問した。富山県知事（左）とWHOの代表者らと表敬訪問した。

富山県は伝統薬の推進と伝統薬の配置サービスで長い歴史と豊かな経験があり、県とWHOの継続的により緊密な協力関係の進展を期待している。と述べ、石井知事は「会議の成果が世界の人の健康増進につながることを期待している」とあいさつ。田宮憲一厚生労働省医政局研究開発振興課長補佐と遠藤俊郎富山大学長も歓迎の言葉を述べた。

富山は伝統薬を推している。今後も富山の強みや特色を生かして、薬用植物の保護に力を入れていく。と述べた。

富山県は伝統薬を推している。今後も富山の強みや特色を生かして、薬用植物の保護に力を入れていく。と述べた。



石井知事（手前左端）と会談するWHO伝統薬部門担当課長（後右から2人目）＝県庁で

記事2 北日本新聞 2011年 10月 15日

薬用植物保護の指針改定へ
富山でWHO専門家会議



WHO専門家会議の開会式であいさつする張奇氏（中央）＝富山市のANAクラウンプラザホテル富山

薬用植物の保護に関するWHO（世界保健機関）専門家会議の開会式が14日、富山市のANAクラウンプラザホテル富山で開かれ、世界各国の専門家らがガイドライン改定を目指し会議の成功を誓い合った。

専門家会議は15日から17日まで、県民会館で開催。21カ国の専門家やWHO本部の職員ら約60人が参加し、WHO

記事3 毎日新聞 2011年 10月 15日

薬用植物保護で結束

富山でWHO専門家会議開会
ガイドライン改訂へ協力



開会のあいさつをする張氏＝富山市内のホテル

薬用植物の保護に関するWHO（世界保健機関）専門家会議の開会式は14日、ANAクラウンプラザホテル富山で行われ、20数カ国の約60人の研究者らが薬用植物保護に関する新ガイドライン改訂へ協力、連携することを確約した。

WHO本部伝統薬部門担当課長の張奇氏が「富山県は伝統薬、家庭配置のサービスで長い伝統があり、この地で緊密な協力的な進展を期待する」と会議の成功を意欲を示した。

石井知事は歓迎のあいさつを述べた。

「会議の成果が人類の健康増進につながることを願う」と述べた。厚生労働省医政局研究開発振興課長補佐の田宮憲一氏、遠藤俊郎富山大学長があいさつした。

会議は15日から3日間にかけて富山市の県民会館で開かれるほか、15日に県民公開講座を開催。関係者は18日に富山大民族学物質館を視察する。

これに先立ち、WHO伝統薬部門の張奇課長らは県庁を訪れ、石井知事と懇談した。

県学生競争展
作品を募集
来月、雨期で開催

記事5 富山新聞 2011年 10月 15日

薬用植物の保護議論

富山 WHO専門家会議開幕

は七月の記者会見では、「という議論の仕方は、オール・オブ・ア・ナツ（全部）という考え、今後シンドではなく、利用も持ち強く、一部伝統的な見込みの時間は残りを模索する考えを表明して、もう一つもいよいよ明している。（相馬敬）

富山 WHO専門家会議開幕

薬用植物の保護に関する世界保健機関（WHO）専門家会議が14日、富山市内のホテルで開かれ、五日間の日程が始まった。

WHOは「薬用植物の保護に関するガイドライン」の改訂を進めており、薬用植物の栽培促進や乱獲防止などを盛り込んだ議論を反映し、改訂指針を取りまとめる。二十一カ国の専門家六十人が十五日



知事と懇談するパッタ二総領事＝県庁で

記事4 北陸中日新聞 2011年 10月 15日

○津波・浜崎、前9・5
急接近フェア（ありそ）
前山田市長書きあけい交
秀（エッセイ）（市議員書
交センター）

十七日、同市新総曲輪の県民会館で話し合
う。十八日は県内の関
連施設を見学する。

県民公開講座もあ
り、十五日午後一時半
から県民会館でWHO
本部科学官の丸山由紀
子氏ら四人が講演す
る。

WHO伝統薬部門の
張奇課長が十四日、石
井知事と懇談し、「薬
用植物の富山で会議を
開催し、長い歴史と豊
かな経験は共有するこ
とに重要性がある」と
話した。（山田晃史）

薬用植物の保護に関するWHO会議

県民公開講座

意外と知らない 身近な 薬用植物



参加無料
事前予約不要

平成23年

日時

10月15日(土)
13:30~16:00

場所

富山県民会館401号室
(富山県富山市新総曲輪4-18)

講演

WHOを身近にー薬用植物を通じて

丸山由紀子(世界保健機関/WHO本部科学官)

身近な薬用植物

佐竹 元吉(富山大学和漢医薬学総合研究所・お茶の水女子大学教授)

漢方薬等の原料となる薬用植物の現状

小松かつ子(富山大学和漢医薬学総合研究所教授)

身近な薬用植物の意外な作用

門脇 真(富山大学和漢医薬学総合研究所教授)

主催：薬用植物の保護に関するWHO会議実行委員会

後援：富山県、富山大学和漢医薬学総合研究所

お問い合わせ

富山大学 和漢医薬学総合研究所 和漢薬製剤開発研究部門 Tel 076-434-7605

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リサイクル推進(A)
この印刷物は、資源の節約に
リサイクルされています。

研究成果の刊行に関する一覧表

雑誌

発表者氏名	論文タイトル名	発表誌名	巻合	ページ	出版年
Zhu S., Kitani Y., Komatsu K.	Exploration of <i>Ephedra</i> resource in Mongolia: From field investigation to molecular identification and chemical evaluation.	J. Trad. Med.,	29	35-40	2012
Ahmed K, Furusawa Y, Tabuchi Y, Emam HF, Piao JL, Hassan MA, Yamamoto T, Kondo T, Kadowaki M.	Chemical inducers of heat shock proteins derived from medicinal plants and cytoprotective genes response.	Int J Hyperthermia.	28	1-8	2012
Marisa Rangel a, Marcia Perez dos Santos Cabrera b, Kohei Kazuma c, Kenji Ando c, Xiaoyu Wang c, Manabu Kato d, Ken- ichi Nihei e, Izaura Yoshico Hirata f, Tyra J. Cross g, Angélica Nunes Garcia a, Eliana L. Faquim- Mauro a, Marcia Regina Franzolin h,	Chemical and biological characterization of four new linear cationic α -helical peptides from the venoms of two solitary eumenine wasps	Toxicon	57	1081- 1092	2011

Hiroyuki					
Fuchino i,					
Kanami					
Mori-					
Yasumoto j,					
Setsuko					
Sekita j,					
Makoto					
Kadowaki c,					
Motoyoshi					
Satake c,					
Katsuhiro					
Konno c,*					

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2
3 **RESEARCH ARTICLE**

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7 **Chemical inducers of heat shock proteins derived from medicinal plants**
8 **and cytoprotective genes response**
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12 KANWAL AHMED^{1,2}, YUKIHIRO FURUSAWA³, YOSHIAKI TABUCHI⁴, HEBA
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23 *(Received 30 June 2011; Revised 11 September 2011; Accepted 23 September 2011)*
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25
26 **Abstract**

27 Environmental stress induces damage that activates an adaptive response in any organism. The cellular stress response is
28 based on the induction of cytoprotective proteins, the so-called stress or heat shock proteins (HSPs). HSPs are known to
29 function as molecular chaperones which are involved in the therapeutic approach of many diseases. Therefore in the current
30 study we searched nontoxic chaperone inducers in chemical compounds isolated from medicinal plants. Screening of 80
31 compounds showed significant Hsp70 up-regulation among them shikonin was most potent. Shikonin was able to induce
32 Hsp70 at 0.1 μM after 3 h without activation of heat shock transcription factor 1 (HSF-1). It also induces significant reactive
33 oxygen species generation. The expression level of genes responsive to shikonin was studied using global-scale microarrays
34 and computational gene expression analysis tools. Significant increase in the nuclear factor erythroid 2-related factor 2
35 (Nrf2, NFE2L2) mediated oxidative stress response was observed that leads to the activation of HSP. The results of gene
36 chip analysis were further confirmed by real-time qPCR assay. In short, the detailed mechanisms of Hsp70 induction by
37 shikonin is not fully understood, Nrf2 and its target genes might be involved in the Hsp70 up-regulation of U937 cells.
38

39 **Keywords:** *Heat shock proteins, Nrf2, oxidative stress, shikonin*
40

41 **Introduction**

42
43 Human exposure to environmental toxicants has
44 been associated with etiology of many diseases
45 including inflammation, cancer, cardiovascular and
46 neurodegenerative disorders. To counteract the detri-
47 mental effects of environmental insults, mamma-
48 lian cells have evolved a hierarchy of sophisticated
49 sensing and signalling mechanisms to turn on or off
50 endogenous antioxidant responses accordingly [1].
51 The ability of cells to counteract stressful conditions,
52 known as cellular adaptive response, requires the

47
48 activation of pro-survival pathways and the produc-
49 tion of molecules with antioxidant, antiapoptotic and
50 proapoptotic activities [2]. Among the cellular path-
51 ways conferring protection against oxidative stress, a
52 key role is played by vitagenes, which include heat
53 shock proteins (HSPs) such as heme oxygenase-1
54 (HMOX1) and Hsp70, as well as thioredoxin/
55 thioredoxin reductase system [3]. Heat shock or
56 stress response is a cellular adaptive response, which
57 contributes to establishing a cytoprotective state in a
58 wide variety of human diseases. When appropriately

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115 activated, cellular stress response has the capability to
116 restore cellular homeostasis and rebalance redox
117 equilibrium [4].

118 Among the many changes in cellular activity and
119 physiology, the most remarkable event in stressed
120 cells is the production of a highly conserved set of
121 proteins, the heat shock or stress proteins
122 (HSPs) [5]. HSPs are also expressed constitutively
123 at normal growth temperatures and play an essential
124 role as molecular chaperones by assisting the correct
125 folding of nascent and stressed accumulated mis-
126 folded proteins, preventing their aggregation [6],
127 assembly/disassembly of multi-subunit oligomers,
128 translocation of proteins across intracellular
129 membranes, process of endocytosis, regulation of
130 apoptosis and cytoskeletal organisation [7]. A
131 number of studies have reported that molecular
132 chaperones can confer cellular and tissue stress
133 tolerance and provide beneficial effects on various
134 pathological states, such as stress ulcers and ischaemia-induced injuries, as well as on diseases associated with protein misfolding and aggregation [8–10].

137 Given the broad cytoprotective properties of the
138 heat shock response, there is now strong interest in
139 discovering and developing pharmacological agents
140 capable of inducing stress responses. Therefore, in
141 the current study we screened some phyto-medicinal
142 compounds for their HSP-inducing activity.

143 **Materials and methods**

144 *Chemical compounds*

148 Chemical compounds used in this study were
149 obtained from the Institute of Natural Medicine,
150 University of Toyama, Japan.

151 *Cell culture*

154 A human lymphoma cell line, U937 was obtained
155 from the Human Sciences Research Resource Bank
156 (Japan Human Sciences Foundation, Tokyo, Japan).
157 The cells were maintained in Roswell Park Memorial
158 Intitute (RPMI) 1640 medium (Sigma, St Louis,
159 MO, USA) with 10% heat-inactivated fetal bovine
160 serum (FBS) (JRH Biosciences Lenexa, KS, USA)
161 and incubated in a CO₂ incubator with 5% CO₂ and
162 95% air at 37°C.

163 *Hyperthermia treatment*

166 Hyperthermia (HT) treatment was used as a positive
167 control for induction of HSPs. Cells were collected
168 and suspended in 2 mL fresh medium in plastic
169 culture tubes, and were exposed to 44°C (±0.05°C)
170 for 15 min in a water bath (NTT-1200, Eyela,
171 Tokyo, Japan). After HT treatment, cells were

172 incubated for desired time at 37°C in humidified
173 air with 5% CO₂.

174 *Assessment of apoptosis*

175 For the detection of early apoptosis and secondary
176 necrosis, Annexin V-FITC kit, purchased from
177 Immunotech (Marseille, France), was utilised
178 according to the manufacturer's recommendations.
179 Briefly, cells were stained simultaneously with
180 propidium iodide (PI) and fluorescein isothiocyanate
181 (FITC)-labelled annexin V and assessed with a flow
182 cytometer (Beckman-Coulter EPICS XLTM).

183 *Determination of cell survival by WST-8 assay*

184 Cell survival was determined using a Cell Counting
185 Kit-8. The cells (2 × 10⁴ /well) in 96-well plates were
186 incubated with various concentrations of shikonin for
187 6 h. After incubation, WST-8 reagents were added to
188 each well. Absorbance at 450 nm was measured
189 using a microplate reader (BioRad, Hercules, CA)
190 after 2 h of incubation with WST-8 reagents.
191 Absorbance is proportionally related to the number
192 of live cells.

193 *Western blot analyses for proteins*

194 Western blot analysis was performed for Hsp70,
195 HSF-1, NRF2 and β-actin by using specific poly-
196 clonal or monoclonal antibodies (Santa Cruz
197 Biotechnology, Santa Cruz, CA) as described in
198 previous papers [11, 12]. For the detection of these
199 specific antibodies the chemiluminescence ECL
200 detection reagents were used following the manu-
201 facturer's instructions (Amersham Biosciences,
202 Buckinghamshire, UK).

203 *RNA isolation*

204 Total RNA was extracted from cells using an
205 RNAeasy Total RNA Extraction kit (Qiagen,
206 Valencia, CA) and treated with Dnase I (RNase-
207 free Dnase kit, Qiagen) for 15 min at room temper-
208 ature to remove genomic DNA.

209 *Assessment of intracellular reactive oxygen species (ROS)*

210 To measure intracellular superoxide (O₂⁻) we used
211 2 μM hydroethidine (HE) (Molecular Probes,
212 Eugene, OR) a dye that is oxidized within the cell
213 and fluoresces when it intercalates into DNA. To
214 measure intracellular peroxides including H₂O₂
215 5 μM dichlorofluorescein diacetate (DCFH-DA)
216 (Molecular Probes) was utilised. DCFH-DA upon
217 entering into the cells de-esterified and then oxidized
218 to dichlorofluorescein (DCF). The fluorescence
219 emission was analysed by flow cytometry [13].

229 *High-density oligonucleotide microarray and compu-*
230 *tational gene expression analysis*

231 Gene expression was analysed using a GeneChip®
232 system with a Human Genome U133-plus 2.0 array
233 (Affymetrix, Santa Clara, CA) spotted with 54,675
234 probe sets, more than twice the molecular probes
235 used in our previous studies [14, 15]. Samples for
236 array hybridisation were prepared as described in the
237 Affymetrix GeneChip® Expression Technical
238 Manual. Briefly, 5 g of total RNA was used to
239 synthesise double-stranded cDNA with a
240 GeneChip® Expression 30-Amplification Reagents
241 One-Cycle cDNA Synthesis kit (Affymetrix). Biotin-
242 labelled cRNA was then synthesised from the cDNA
243 using GeneChip® Expression 3'-Amplification
244 Reagents for IVT labelling (Affymetrix). After frag-
245 mentation, the biotinylated cRNA was hybridised to
246 the GeneChip array at 45°C for 16 h. The arrays were
247 washed, stained with streptavidin-phycoerythrin, and
248 scanned using a probe array scanner. The scanned
249 chip was analysed using GeneChip Analysis Suite
250 Software (Affymetrix). The obtained hybridisation
251 intensity data were converted into a presence or an
252 absence call for each gene, and changes in gene
253 expression level between experiments were detected
254 by comparative analysis. The data were further
255 analysed using Gene-Spring software (Silicon
256 Genetics, Redwood City, CA) to extract the signif-
257 icant genes [16, 17]. To examine gene ontology,
258 including biological processes, cellular components,
259 molecular functions, and genetic networks, the
260 obtained data were analysed using Ingenuity
261 Pathways Analysis tools (Ingenuity Systems,
262 Mountain View, CA), a web-delivered application
263 that enables the identification, visualisation and
264 exploration of molecular interaction networks in
265 gene expression data. The gene lists identified by
266 GeneSpring containing Affymetrix gene ID and the
267 natural logarithm of normalised expression signal
268

286 ratios from GeneChip CEL files were uploaded into
287 the Ingenuity Pathways Analysis system. Each gene
288 identifier was mapped to its corresponding gene
289 object in the Ingenuity Pathways Knowledge
290 Base. These so-called focus genes were then used
291 as a starting point for generating biological net-
292 works [16, 17].
293

294 *Real-time quantitative PCR assay*

295 Real-time quantitative PCR (qPCR) assay was per-
296 formed on a real-time PCR system (Mx3000P,
297 Stratagene, Tokyo, Japan) using SYBR PreMix
298 ExTaq (Takara Bio, Shiga, Japan) or Premix
299 ExTaq (for the use of TaqMan probes; Takara Bio)
300 in accordance with the manufacturer's protocols.
301 Reverse transcriptase reaction (Omniscrypt Reverse
302 Transcriptase, Qiagen) was carried out with DNase-
303 treated total RNA using an oligo (dT) 16 primer.
304 Real-time qPCR assay was performed using the
305 specific primers listed in Table I. Each mRNA
306 expression level was normalized to the mRNA
307 expression level of GAPDH.
308

309 **Results**310 *Screening of medicinal compounds for Hsp70 up-*
311 *regulation*

312 Eighty medicinal compounds shown in Table 1 were
313 examined for their ability to induce Hsp70 up-
314 regulation. The screening procedure was accom-
315 plished by treating U937 cells with the compounds 1
316 to 10 at 0.1 mM for 24 h followed by western blot
317 analysis. Among the tested samples no band of
318 Hsp70 was observed with compounds 3–5 suggesting
319 the toxicity of the dose (data not shown). Therefore,
320 we selected compounds 3–5 and did concentration-
321 dependent screening for Hsp70 up-regulation. U937
322 cells were treated with 0.1 μM, 1 μM and 10 μM for
323
324
325
326

327 Table I. List of 80 chemical compounds that were examined for their Hsp70 inducing ability. 328

272	1. Aconitine	17. Bufotalin	33. Epihesperidin	49. Gomisin N	65. Paeoniflorin	329
273	2. Albiflorin	18. Capillarisin	34. Ergosterol	50. Hesperidin	66. Paeonol	330
274	3. Alisol A	19. Capsaicin	35. beta-Eudesmol	51. Hirsutine	67. Palmatine chloride	331
275	4. Alisol B	20. Catalpol	36. (E)-Ferulic acid	52. Honokiol	68. (S)-Perillaldehyde	332
276	5. Alkannin	21. (E)-Cinnamic acid	37. Geniposide	53. Hypaconitine	69. Puerarin	333
277	6. Amygdalin	22. Cinobufagin	38. Geniposidic acid	54. Icarin	70. Rhynchophylline	334
278	7. Arbutin	23. Cinobufotalin	39. Gentiopicroside	55. Isofraxidine	71. Saikosaponin a	335
279	8. Astragaloside IV	24. Coptisine chloride	40. [6]-Gingerol	56. (Z)-Ligustilide	72. Saikosaponin b2	336
280	9. Atractylenolide III	25. Corydaline	41. Ginsenoside Rb1	57. Limonin	73. Saikosaponin c	337
281	10. Aucubin	26. Curcumin	42. Ginsenoside Rc	58. Loganin	74. Schizandrin	338
282	11. Baicalin	27. Dehydrocorydaline nitrate	43. Ginsenoside Rd	59. Magnolol	75. Sennoside A	339
283	12. Baicalin	28. Dehydrocostuslactone	44. Ginsenoside Re	60. Mesaconitine	76. Shikonin	340
284	13. Barbaloin	29. Dihydrocapsaicin	45. Ginsenoside Rg1	61. Naringin	77. [6]-Shogaol	341
285	14. Berberine chloride	30. Dimethylesculetin	46. Glabridin	62. Nodakenin	78. Sinomenine	342
	15. Bergenin	31. Eleutheroside B	47. Glycyrrhizic acid	63. Osthole	79. Swertiamarin	
	16. Bufalin	32. (-)-Epigallocatechin gallate	48. Gomisin A	64. Oxymatrine	80. Wogonin	

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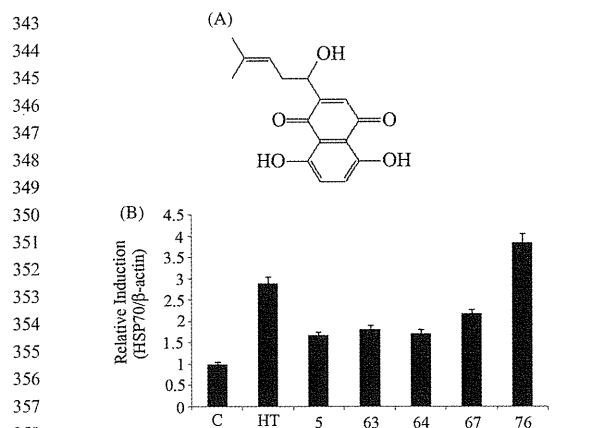


Figure 1. (A) Chemical structure of shikonin. (B) Bands of Hsp70 induced by five active compounds at $1\mu\text{M}$ concentration were quantified by densitometry and normalised with β -actin. Hyperthermia (HT) 44°C for 15 min was used as a positive control. Bars indicate standard deviation ($n=3$).

24 h followed by western blot analysis. Significant Hsp70 up-regulation was observed with compound 5 at $1\mu\text{M}$ (densitometric ratio: control: 1.0 ± 0.08 , $0.1\mu\text{M}$: 1.25 ± 0.12 , $1\mu\text{M}$: 1.50 ± 0.14 , $10\mu\text{M}$: 1.12 ± 0.13 , mean \pm SD, $n=3$) while compounds 3 and 4 did not show any significant increase at all concentrations as compared to control. From these results we selected $1\mu\text{M}$ concentration for the screening of medicinal compounds.

Out of the 80 compounds, five (5, 63, 64, 67 and 76) showed significant Hsp70 up-regulation. Among them compound 76 (shikonin) was the most potent (Figure 1A, 1B).

Effects of shikonin on apoptosis induction in U937 cells

To determine the non-toxic concentration of shikonin, cells were exposed to the drug concentration-dependently for 6 h. This was followed by measurement of early apoptosis and secondary necrosis by annexin V FITC/PI staining using flow cytometry. No apoptosis was observed at $0.01\mu\text{M}$ and $0.1\mu\text{M}$, significant apoptosis was observed at $1\mu\text{M}$ concentration (Figure 2A).

Effects of shikonin on cell survival

To further confirm the non-toxic concentration, cell survival assay was performed with 0.01 , 0.1 and $1\mu\text{M}$ concentrations after 6 h incubation. The results also showed the toxicity at $1\mu\text{M}$ while no toxicity was observed with 0.01 and $0.1\mu\text{M}$ concentrations. (Figure 2B).

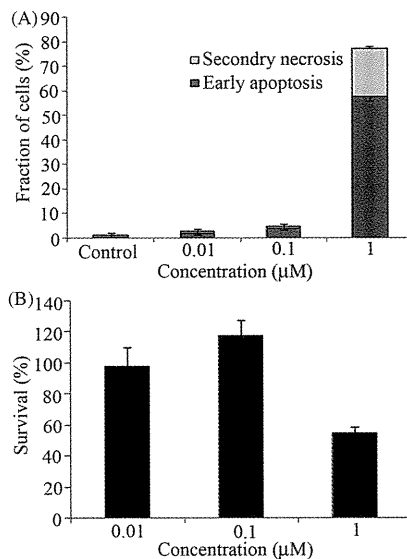


Figure 2. (A) U937 cells were treated with shikonin concentration-dependently for 6 h followed by assessment of early apoptosis (black bar) and secondary necrosis (grey bar) by flow cytometry. Bars indicate standard deviation ($n=3$). (B) Cell survival assay was performed with 0.01 , 0.1 and $1\mu\text{M}$ concentrations of shikonin. After 6 h incubation WST-8 reagent was added to each well. Bars indicate standard deviation ($n=3$).

Effects of shikonin on ROS formation

Previously it has been reported that shikonin can induce ROS formation [18]. Therefore, in the current study we investigated that whether at non-toxic concentration shikonin can induce ROS. Cells were exposed to $0.1\mu\text{M}$ shikonin and the levels of DCF and HE fluorescence were monitored time dependently via flow cytometry. Intracellular peroxide level was increased as early as 30 min after treatment (Figure 3A), while no intracellular O_2^- was observed (data not shown).

Time-dependent effects of shikonin on Hsp70 induction

U937 cells were treated with $0.01\mu\text{M}$ and $0.1\mu\text{M}$ shikonin time-dependently, followed by western blot analysis. Up-regulation of Hsp70 was observed with $0.1\mu\text{M}$ after 3 h incubation and continued to increase time-dependently, while no up-regulation of Hsp70 was observed at $0.01\mu\text{M}$ shikonin (Figure 3B, 3C).

The transcription of HSP genes is regulated by transcription factor HSF, which senses cellular exposure to stress and turns on rapid induction of HSPs [19]. Therefore we examined the effects of shikonin on the activation of HSF1. U937 cells were